# Automatic Construction of UMLS Metathesaurus with Deep Learning

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## Unified Medical Language System (UMLS) Metathesaurus

Started in 1986 by the National Library of Medicine (NLM)

#### Overcome barriers to effective retrieval of machine-readable information

- The variety of ways the same concepts are expressed by different terminologies (MeSH, MedDRA, RxNORM, ICD-10, SNOMED CT, etc)

- ~ 10 million English medical terms
- From 210 source vocabularies
  - General
    - Anatomy (FMA, Neuronames), drugs (RxNorm, ATC, First DataBank), medical devices (UMD, SPN),
       clinical terms (SNOMED CT), information sciences (MeSH), administrative terminologies (ICD-9/10)
  - Specialized
    - Nursing (NIC), psychiatry (DSM, APA), adverse reactions (MedDRA)
- Grouped into ~ 3.85 million concepts

Used in areas such as patient care, clinical coding, information retrieval, knowledge exploration, and data mining

#### Integrating Subdomains

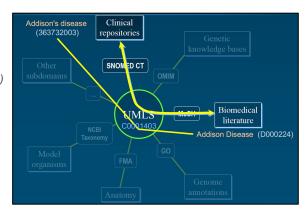
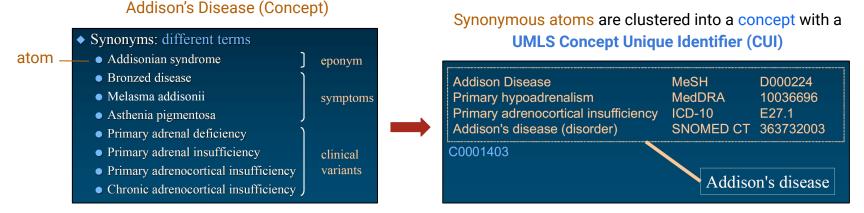


Image from **Unified Medical Language System Overview**by Olivier Bodenreider

## Unified Medical Language System (UMLS)

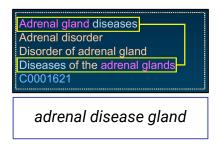


Images from **Unified Medical Language System Overview**by Olivier Bodenreider

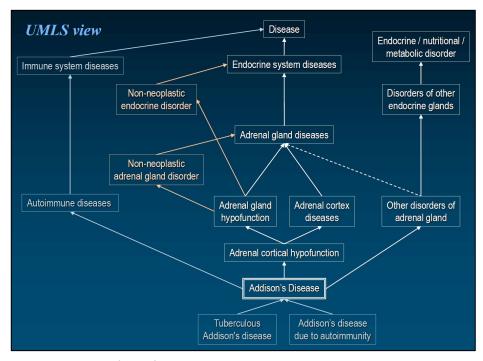
## Construction of UMLS Metathesaurus (Updates bi-annually)

#### Lexical Knowledge

Lexical Variant Generator (LVG)



- Semantic Pre-processing
- UMLS Human Editors



Images from Unified Medical Language System Overview by Olivier Bodenreider

## **Motivation**

The current approach in adding new resources from identifying lexical variants to manual audits can be both arduous and time-consuming.

(~ 10 million English medical terms, ~ 3.85 million concepts)

## Objectives

The project explores the realm of supervised machine learning approach (Deep Learning) to

- 1. Identify **synonymy** and **non-synonymy** among UMLS concepts at the atom level
  - Given two atoms, are they synonymous (same CUI)?
- 2. Investigate Deep Learning approach could emulate the current building process

## **Problem Formulation**

#### Approach 1 (Classification task):

- Training Data: ~ 10M English language atoms and each with its own CUI assignment
- We can train a classification model to predict which CUI should be assigned to a given "new" atom (since atoms having the same CUI are synonymous).
- Input: Atom -> Output label: CUI
- Challenge: ~ 3.85M softmax outputs (extreme classification task)

#### **Approach 2 (Similarity task):**

Learn similarities between atoms within a CUI and dissimilarities between atoms from different CUIs.

A fully-trained model should identify and learn scenarios where

**Lung** disease and disorder

Two atoms that are *lexically* similar in nature but are not synonymous

**Head** disease and disorder

Addison's disease

Two atoms that are **lexically dissimilar** but are **synonymous** 

Primary adrenal deficiency

### Traditional Neural Network Architecture

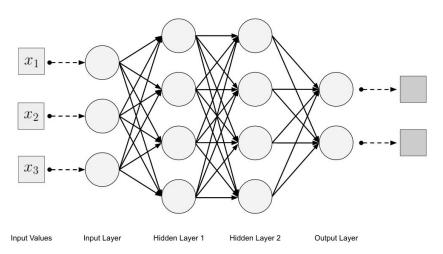
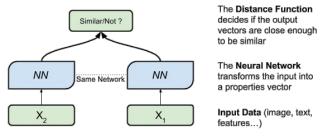


Image source: https://www.oreilly.com/library/view/deep-learning/9781491924570/ch04.html

Feedforward Neural Network (Multilayer Perceptron):

Not suited for Pairwise-similarity task

## Siamese (Twin) Neural Network



#### Image source:

https://aws.amazon.com/blogs/machine-learning/combining-deep-learning-networks-ganand-siamese-to-generate-high-quality-life-like-images/

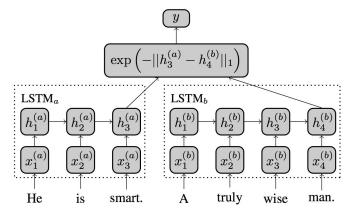
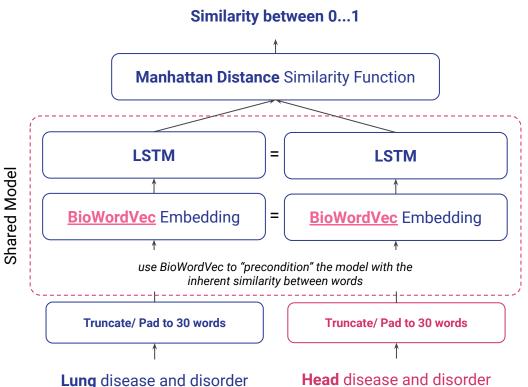


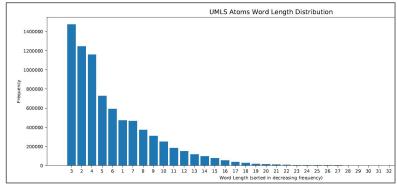
Image source: Siamese Recurrent Architectures for Learning Sentence Similarity 7

## Siamese-LSTM





Zhang Y, Chen Q, Yang Z, Lin H, Lu Z. BioWordVec, improving biomedical word embeddings with subword information and MeSH. Scientific Data. 2019.



#### UMLS Atoms Word Length Distribution

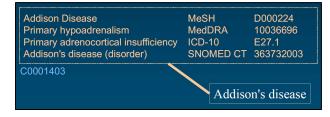
(Word length 30 covers 97% of atoms in the UMLS)

## Dataset (2019-AA UMLS) and Feature Engineering

#### **Positive Pairs (Synonyms)**

• (CUI)-asserted synonymy between atoms (~15 million pairs)





#### **Negative Pairs (Non-synonyms)**

Ideally, we want to generate all negative pairs (1 atom against atoms from other non-related CUIs)
 (~ 10 million atoms \* 10 million atoms)

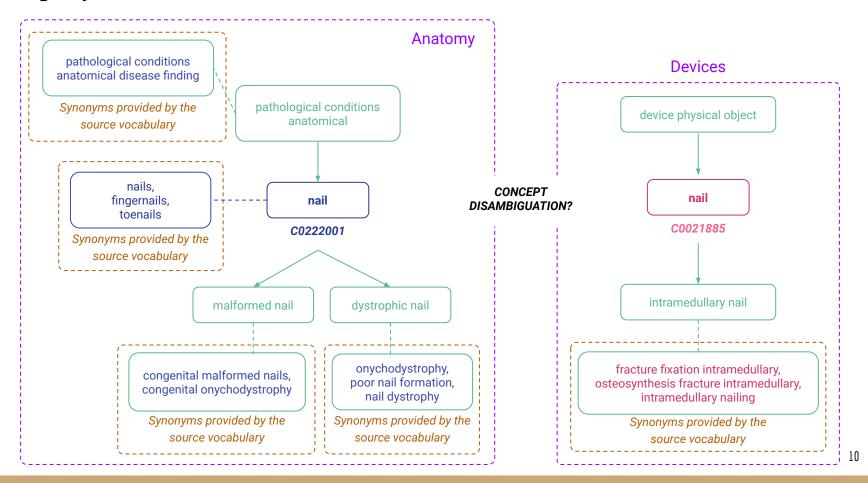
Class Imbalance: Number of *Non-synonyms* > Number of *Synonyms* 

Intuition: What we want are interesting negative pairs that are lexically similar but differ in semantics.

• Heuristic Approach: Use **Jaccard Index** to generate **negative pairs** for atoms with **high Jaccard Similarity** (Sort and filter top  $\sim$ 15 million pairs)



## Going beyond atoms... Let's Contextualize!



#### 2. "Base" (Atom lexical features)

- + Synonyms provided by the source vocabulary
- 3. "Base" (Atom lexical features)
- + Hierarchical-Context(atom)
- + Semantic Group

1. "Base" experiment

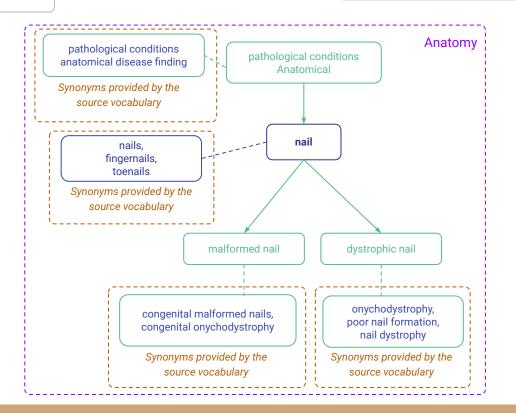
(Atom lexical features)

- 4. "Base" (Atom lexical features)
- + Synonyms provided by the source vocabulary
- + Hierarchical-Context(atom)
- + Semantic Group

#### 5. "Base" (Atom lexical features)

- + Synonyms provided by the source vocabulary
- + Hierarchical-Context(atom)
- + Synonyms of the Hierarchical-Context(atom)
- + Semantic Group

## Experimental Setup



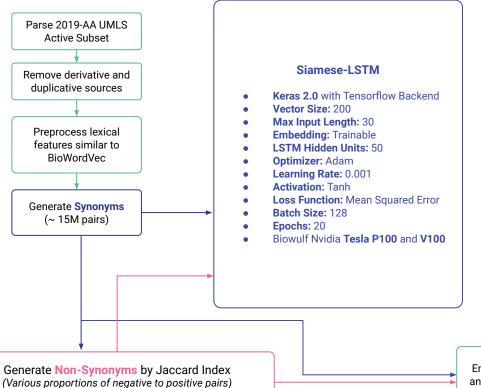
#### Architecture Fully Connected Layer (50) Fully Connected Layer (128) Similar/Not? Concatenated Vectors "Base" Vector "Context" Vector "SG" Vector NN NN Same Network LSTM **LSTM LSTM** Learn word order and features Learn word features/ context Learn word order and features CNN Conv1D Anatomy pathological Extract word features conditions anatomical disease pathological finding conditions BioWordVec anatomical BioWordVec BioWordVec 30 nails, 30 **Embeddings Embeddings Embeddings** fingernails, toenails nail anatomical conditions dystrophic congenital malformed nail disease anatomy nail nail dystrophic dystrophy finding onychodystrophy, fingernails congenital malformed nails, formation poor nail formation. congenital onychodystrophy Extract only unique lexical features to malformed nail dystrophy nails enrich the "base" and sort to Context Bag onychodystrophy "eliminate" word order randomness pathological 17 toenails Pontes, E. L., Huet, S., Linhares, A. C., & Torres-Moreno, J. M. (2018). Predicting the Semantic

Textual Similarity with Siamese CNN and LSTM. arXiv preprint arXiv:1810.10641.

## Methodology Overview

Experiment 1 (5-fold Cross Validations)

Experiment 2, 3, 4, 5 (5-fold Cross Validations)



#### Siamese-CNN-LSTM

- Keras 2.0 with Tensorflow Backend
- Vector Size: 200
- Max "Base" Input Length: 30
- Max "Context" Input Length: 60
- **Embedding:** Trainable
- LSTM Hidden Units: 50
- LSTM Activation: Tanh
- CNN Filters: 100
- Window Size: 5
- CNN Activation: ReLU with Batch
  Normalization
- Fully Connected Layer 1: 128 with ReLU
- Fully Connected Layer 2: 50 with ReLU
- **Optimizer:** Adam
- Learning Rate: 0.001
- Loss Function: Mean Squared Error
- Batch Size: 128
- Epochs: 20
- Biowulf Nvidia Tesla P100 and V100

Enrich with "Context" and "Semantic Group"

## Results & Evaluations (based on optimal runs)

Model/ Performance Metrics	Base	Base	Base	Base	Base
		+ Source Synonymy	+ Hier. Context + Semantic Group	+ Source Synonymy + Hier. Context + Semantic Group	+ Source Synonymy + Hier. Context + Hier. Source Synonymy + Semantic Group
Accuracy	0.9333	0.8720	0.9486	0.9520	0.9541
Precision	0.7828	0.8654	0.7643	0.8296	0.8009
Recall	0.7379	0.8874	0.8381	0.9038	0.8978
F1-Score	0.7597	0.8763	0.7995	0.8428	0.8466
Matthew CC	0.7214	0.7441	0.7712	0.8173	0.8215
Specificity	0.9659	0.8560	0.9640	0.9601	0.9633
Sensitivity	0.7379	0.8874	0.8381	0.9038	0.8978
False Positive Rate	0.0341	0.1440	0.0360	0.0399	0.0367

#### Observations:

- Source synonymy is responsible for achieving high precision and overall F-1 score.
- Adding hierarchical context trades precision for higher recall.
- Adding source synonymy, hierarchical context, and semantic group give an overall boost to accuracy and recall.
- However, adding source synonymy of hierarchical context did not yield any noticeable improvement.

## Examples of True Positives and True Negatives Correctly Identified

True Positives (Synonyms) Correctly Identified				
nail clipper	cutters nail			
injury of salivary gland	salivary gland injury			
avulsion	fracture sprain			
True Negatives (Non-synonyms) Correctly Identified				
fingernail	infection of fingernail			
product containing only iron medicinal product	product containing only levorphanol medicinal product			
medical and surgical gastrointestinal system insertion ileum via natural or artificial opening endoscopic infusion device	medical and surgical gastrointestinal system revision <b>stomach</b> via natural or artificial opening endoscopic <b>other</b> device			

## Examples of False Positives Identified and False Negatives Not Identified

False Positives (Non-synonyms) Identified				
finding of wrist joint	finding of knee joint			
malignant neoplasm of upper limb	malignant neoplasm of muscle of upper limb			
skin wound of axillary fold	skin <b>cyst</b> of axillary fold			
False Negatives (Synonyms) Not Identified				
hla antigen	human leukocyte antigen			
pyelotomy	incision of renal pelvis treatment			
routine cervical smear	screening for malignant neoplasm of cervix			

## Conclusion & Future Work

- Deep learning approach provides good performance in identifying synonymy among atoms.
- Adding source synonymy yields better precision and overall F-1 score.
- Adding hierarchical context trades precision for higher recall.
- Adding source synonymy, hierarchical context, and semantic group give an overall boost to accuracy and recall.
- **Limitation:** This approach does not address the *inter-concept* and *semantic type categorizations* (other components in the UMLS Metathesaurus).
- Future work: How can the models be used in conjunction to complement the current lexical processing and human editors.

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## Thank you!